30

SEQUENCE LISTING

SEQ ID NO: 1 (Mouse GCR1/Fragilis Nucleic Acid)

Mouse GCR1 (Fragilis) full length nucleotide sequence

SEQ ID NO: 2 (Mouse GCR1/Fragilis Amino Acid)

Mouse GCR1 (Fragilis) amino acid sequence

MNHTSQAFITAASGGQPPNYERIKEEYEVAEMGAPHGSASVRTTVINMPREVSVPDHVVWSLFNT LFMNFCCLGFIAYAYSVKSRDRKMVGDVTGAQAYASTAKCLNISTLVLSILMVVITIVSVIIIVL NAQNLHT

SEQ ID NO: 3 (MOUSE GCR2/STELLA NUCLEIC ACID)

20 Mouse GCR2 (Stella) full length nucleotide sequence

SEQ ID NO: 4 (MOUSE GCR2/STELLA AMINO ACID)

35 Mouse GCR2 (Stella) amino acid sequence

MEEPSEKVDPMKDPETPQKKDEEDALDDTDVLQPETLVKVMKKLTLNPGVKRSARRRSLRNRIAA VPVENKSEKIRREVQSAFPKRRVRTLLSVLKDPIAKMRRLVRIEQRQKRLEGNEFERDSEPFRCL CTFCHYQRWDPSENAKIGKN

SEQ ID NO: 5 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; similar intron-exon structure as mouse-Stella. AC094826 contig No.5 (22671 - 27595: contig of 4925 bp in length)

ACACGAAGCGGACTCCCCGCATCATTCACGTAGACCCGCCTTCTGCTTTCCCTGTCGGGGTTTTG 10 GGAAGCCCGGCGCCCTCTCTCTCACCTTGCTCCACTAGCACGCGGCTGTTTTCACTGAGCCCA GCACTGGCTAAGTGGAGCACCAGGAGTTTCAGGCTATCCTTCAGAGGGCAAGGTGTAGTCCATGG TGGGCTACAGGAGACCCTCTCTCTCCGTGAGTACAGAGGGCAAACCCAAGCCAGACAGGGGTGA TGATTAGGAACATACCTTCGTCGGGGAGAAAATACCGGTTCATATAGGAATAAGAGGAACCAGGA GGTAGTTAAGGCTGTGGTGTCTGGTTGCGGGGGTTTTTTGACTCTCAACAACCACGTTCAGAACGTG 15 CTGAGTTTTTATGATGGTGTAGAATTTCCTTATCAGCAATTGGTCTCCGCGGTGTTTCTTTTCT TTTTTAATTTTTAAGTATAATTTGGTGTTTGAAGCAACTGTACTTGGACTAGAACTCCCTGTGT TTGTTTTGATTACGTTGTAGCCCAGGCTGGGCTCAATCTCAATCCTCCTGCCTCTGCCTTCTAAA 20 ATTTTGGCTCTTTTTTTTGGAGCTGGGGACCGAACCGAGGGCCTTGTGCTTCCTAGGCAAGCGC TCTACCACTGAGCTAAATCCCCAACCCCAGTGTAGCTTTATTTTTAAGAACAGGAGTCTTGTTTC TCAAAACAGTTTCTCTGTAGCCCTGGTTGTCCTGGAACTCCGTAAACCAGGCTGGTTTGGGACTC TGCCTTTAAAACACTGGGACTAAAGGCGGTACCACCTCCGTGGGCTACACCGGAATCTTTTAAGC TTCATTTGAACCGGGGCTTTTTCTTTTTCTCACCCACTTTCTGGAAGCGATTTTCCTGCTAAATT 25 TCCATTCCTGGTAAATGACTCTGAGGGGAAATAGGAACCCAGAATAGATTGAGCCGGGGGCTACC TGGGACCCCGCACCCCCCAGCCGCTGTTGAAGCTCTTTGCCTGAGGGGCCTCCGGGTT TGATACCTCCTAGCACTCCGGGCTGAGGGCGTGGCTCGGGAGGAGCCATTCCTTTGGAGAGGAAA ACAACTGCTGGCCTTGAATCTGCCCTAATACCTGACAGTTACATGGGACCTCCTTATTTCCACAG GATTCTTTAGTCTTTGGGAGATTTTCAAATCTTGAGACTGCTCAACCCTTCCTGGCCTAAC 30 AGCTGGCTCACCCTTGGTGTCACTTTGCTTTAACATTCGGAAAAGTTGTGGTAAGTTTCCTGTAT TTACATAGATGTCCGGAAGCATTGGAGCAGGTCAATTAGATTTAGGTGGAAACAGCCTGTTTTTTG GAAAGCTTTCCAGGGCGAAAATGAACCCAGAGGCACTATTGGGCAAGCCCTCCGGCTAAGCAAC 35 ACAATTGGCTGCAGGGGTCTCTGGAAGAGGTGTGAGACAAGAGAGAATATGCAGGTTTCAGGACC TCTGAACTAGAGTTAGGCTGTAACATTGTAACATTGCTGTAAGCAGAACAGCCCATGGTAAG AAGCTCAGTGGATCTCTACAAACACTAGGATATCTGCTCAGGGTTTATGACCAGGCCCTGTGCAT ATGGTTTGCTTCTTGTTGGCCCCCTCTCTTGAAGAGGGGGTGATTATCTGTTACCCACTTCCTTGTT TCTCTGGGGTATTACCTTGCAAAATGCAAAATGATATACTTCACTAATGTCTCCATCTTCTGTTT 40 CAGAAATCCTACAACCAGAAACACTAGTAAAGGTCATGAAAAAGCTAACCCTGAACCCCAGTGCC CAGAAGTGAAAGAATCATGAGGGAAGTTCAAAGCGCCTTTCCCAGGAGAAGGGTCCGCACTCTGT TGTCCGTGCTGAAAGACCCCATAGCAAGGATGAGAAGATTTGTTCGGGTGAGTTGCGTTTGTGGG CGGGGCATAGATCTAAGAGCAACTCTAGCCTCAGGAATGGCACCTAGGTTAAACAGGGAATGTAG 45 ACAAGGATAGTGACTACCTGTGATTCCCAGCTCAAGAAAACAAGCTCCAAGGCTATCCTCTACTG TTTCAGACTCCCCCATAGTCCAAACTGGCCCTCCAGTTCAGTCCACGGTCCTGCTTCTTCCC CGGTGCTAGGCTTTTGAGTGATAAGGCTGACTTAGACTGGATCTCAGAGCTGAAGTGGACCTGTT AGTCTTTGTAGACCAGGCTGGGGTGGTTTCTGCTTTCTCAGCGCCCTAGCTCACATAGTAGGCATT

TTAACTTTGTCTTAATAGTAATTTGAGTAATTTTGTTTTTTCTCTTTGAAGATTGAGCAGAGACAAA TCTTGTTTTTACTGTTTCCTTAGACAAGGAGTGTGTATGTGGAGAGTTACCTTCTCAACACAGGG GAGCCATTCAGATGTCTCTGCACTTTCTGCCATTATCAGAGATGGGATCCTTCTGAGAATGCTAA AATCGGGCAGAACCAGAAGAATTAGGGCAGTTTGAATTGTACACCGTCCTTGCCGTTAACGGTGC CATGCAGCAGATGTGAAAGCTGTTTTTTTTTTTTAAGATTAAACTTTTCTTGGTGCTGGGGAAATC TCTTCTAATTGCTAACCTTTAAATTATATAGGATGTGTGACATTTGGATTCATGGGAATGACAGA 10 TTTACCCAAGAATTGAGCATGAGTCAAAGCCTGGTAGTTTGATTTAGAAGGTAATTGGAATAAAT CTTTTTATTTTAGATTTTCTAGTTTGCAGAGAAATTTGTAATAAAGGCAAATTTGTTATCTTTAA TAAATACAGAACAGATTAGAATGAGCCATTGGAGATGGGGGACTCGTTTTTTACAGGTGCATGTG TGGGTGTGTGATGTTCAGAGTTCAATGTGTGCTACCCTGTATTTCTGCTTGAGGCAAGGTCTCCA TGAGGCCTAGCTGGTCTAACTCCTGGTCCTGCCTTTTGTTTTCCCCCTGAGTTTTGACACCATAGG 15 CTTGTCGGCAAGATCTGGAAGAGGCTTGATGTTTTGTGTTTTGTGCTGTGTAATAAACAATTGGTTG ATAATTGTATGCTTTATTTCCTGAGAGAAGTGTCAGGAAAGGAGGAGTTAGGAAGAAGCCCCAG GCTGGGGTTAAGAGCACTGGCTGCTTTTCCAGAGGTCCTGAGTTCAATTCCCAGCAATCACCTGG TGGCTCCCGAACATCTGTAACAGGATCCAATGCCCTCTTTTTGGTGTGTCTAAGAACTCCCTAGGC 20 GGAACCGAACCCAGGGCCTTGCGCTTGCTAAGCAAGCGCTCTACCACTGAGCTAAATCCCCAACC CCTACAATGGCCTTTTTCTACCTGCTTTTGAATTATCAATAAAAGACTGGGGCAAAAGAAGGCT 25 AGAGAGAATGTGAGGTGTGTATGAAGATTGTGTGTGGGGTTGGGGGATTTAGCTCAGTGGTAGAGT CAAGGTGTGTATCAAGAGTGTGTGTGAGAGTGAAAGGGTAATGAACAGAGGTGTGCATGAGCGTG GGAGTTTGAGAAAAGAAAACAGCAATAAAAAAAAAAAGCAGAGTGCACGAGAGAATGCAGAGTGTG 30 TGCTTCCAGTGGAGAACTCTGATTCTATGTTGAGGCTGGACCCTGGCAATAGTGGGCTTCTTGAA AAATAGTCAAAGGAAACAGTGCTACACCATGGACTTAAGCCTTTAGACTCAGTTCTGGCTTCAAG AGCAGCTGTCAGAAAATAAGTGATGAACTACTTGCAGTCGAACTCGAATC

35

SEQ ID NO: 6 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC097234 (131006 132449: contig of 1444 bp in length)

SEQ ID NO: 7 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC093991 (1 - 7657: contig of 7657 bp in length)

CCTCCAAAAGAGTGGAACACTTCAACTGCCAGATCCAAGATACTGAAATGGGTAGCATGCTGGAG AAAGAATTCAAAAGTTAGGTAGAGAATCTGGTTGAGCAGAGCACTTGCTTTTCTTCCAGAGGATC TGAGTTCAAGTCCCAGGACCTATATCACAGTTTTCTGTAACTCTAGCTCCAGAGGGTCTGACACT 20 TCTGTTCACTGTGGGCACCTGCATTCACAGACAAACATAAAGTAGTTCATCACCCCTTTTCACAGA AAACCCACAGCATGTGAGGAAATCCGGGTCTCTGCGCAATGCCCCCACAGCAGAAGGGGGGGAGCT GGAGAGATGGTTCATCTGTTAGCCCATTTATTGCTCTTGAAGAGAACCCAGGGTCATCCATAGCA CCCATAGCAGCTCACAACCATCTCCAGTTCCAGGAGATCCAATGCCCTGTTGTGACCTCAGGTAC 25 CAGGCATACACAATGAACCTGCACACATACAAAAGTCCATAGAGCCATAGTTACCATTGTGAGCT CTGAGAACCAAATCCGTGTTCTCTGCAAGAGCGACATGCACGCTGAGAACCAGGCACCTTTCCCA $\tt CTGCCTCTTGAGACAAGATCTCACTATGTAGTTCACACTGGCTTCCGACTTGCCACCATCCTCCT$ GCCTCTGCCTATAAAGAATGCTAGGATTATATAGGTACAAAATCACACCTGGCTGTTAAGGTTTT TCTGGCTGTTTTTTTTTCACCCCCATGAATGATTTTGAAAATAGTTGAGCTGTTTACATTAATA 30 AAACAAAATCAGATGGAGACTATATGTCATTATTCATGAATCAAATGACTAGTAACAATACTGAG TAGTTTTGCTTTTGTTTTTGAGCAGGCTCTCACTGTGTAGTCCTGGGTGATCTGGAACTTAC 35 GTGTGTGTGTTCCCCGGAGGCCATGTAGGCGCATGCTTGAACCAGAACCAGAGGAAGTGTGTT TACAGTTACCCTGGGAGGCCAGAAGAGGGCCAGGAGATGCCCTGGAACTGGAATTTCTGGTAGTGG TTAACTGCCTAAAGTGCTGGGACCTAACACTCTTAACTTCTGAGCCATGGCTCTAGTCCTGGGGT 40 CTCATGGTGACACAATTGAGCATTGAGAGCAGCTACAGACCGATTAGATCAGACTTATTAAATTC TTGCCAAGTATGTGGTGACGCAGGCCTGCAATGCCAGTAACTTTGGAGACTGAGCCAAGCAGATC ACCTGAGCCTAGAGACTCAAGGCCACCCTGGACAACATAGAGATATCCTGTTTCAAAATGAAACA AGCTAAGTTCTTTGTACATAGCAGCCTCTCTATTGACTGTGGCAGGGCAGCTGACAGTGTTCTCA 45 CCTAGTCACAGATGTTCTTTCTAGAGGGAACAGACCCGATGAATACAAACATTTTTAGCTCAAGT AAAAGTCTATACTATGAAGGAACTACTTCTTCAAACATCATAACATTTAAAAATGAGAGATTTTAC AAACCTTTTTTTAAAGATTTATTTGTTTATGATAAGTACACTGTCACTGTCTTCAGACACACCAG AATTGGGCATCAGATCTCATTACAGATGGTTGTGAGCCACCATGTGGTTGTTGGGAATTGAACTC 50 CTGGGGACCGAACCCAGGGCCTTGTGCTTGCTAGGCAAGCGCTCTACCACTGAGCTAAATCCCCA

ACCCCCAGCCAGTGCTCTTAACTGCTGAGCCATCTTCCCAGCCCCAACATCAATTTTTTGGTCTAG ATGTTTTACCCTGGTGCCATGCCATCTCGATGGCCCTTGTGGCAGGGGTGCCGGTAAGGCAG CCCCTAGGGCATGAGTTAGGGAGAGCAAAACCTGACCCAGAACCTGACTGCCATGAAGTGATGGA ACTTGACACATGCTACAGTCATCTGAGAGAGAGACTTAATTGAGAAAATGCCTCTGTATTTTCTC 5 GATTGGTATTAGAAGTAGAATATTGCTGTAACAGACCCTAACCATGTTCTCTTGGGGAGGATTGT GGGAAGACTTTGGAACTTGGAACAGGAGAAGCCATTGGGTACTTAGAGCTTAATGGGC TGTTCTGTGGAGAAGGTGCTGGAGAAATGCGGATGATACTTGTAAAGTTTGAGAGCACC ${\tt TCAAAGATGTTCAGGACAGTGTGTGCAATACATTTGAGTTAAGAATCTATGGTGTCTGGTCAGCT}$ 10 GGAGCTGAAGATTCAGCTGTGATTAATAAGACCACTAAAGTAAAACTTTTGCTTTACTGGTACAA TCAGTGCTGGTTAGCTAAGGGTTGACAGATGAGCAGTGACTAATAAGAGACTGGCATCAGAAACT GATCCAGAGAGAGCCAAGGCTGCATCTCAAACTGGCAGCCAAATTTGATCACATGTAAGAATCTC CCTCATGGGGGTTGGGGATTTAGCTCAGTGGTAGAGCGCTTGCCTAGGAAGCACAAGGTCCTGGG 15 CAGGCTTTGGTGGCATGAGAGCTTTAGGGTTGAAGGATCATGGAGAGCAGCCGAGGCTCCGCACC ATGTGGCGGGCAGAGCCCAGTTACCACAGAGACACCAGCATATTTGGAGGTGCCAGGA TCATGGATAATTGCCTAAGACAGGAGGCTGGCCTGACTTTGTAGGACAAGCTCCATGATCTGTTT GGCAGGACTGGAGAAACAGAGCTGTAAGGGAAAATGAGGACACAGCTGTTCCAAGATATGATTGG AGAGAAGGGTTTCATTGCAGATCTGAGGAAGAGGACAGCCAGAGAGGCCATCTGGAAGGGTCCAGA 20 TTGAACTGGGTCATGAGAGAGAGAGGGCTAAGAGAGCCAAAAGAGCCTGTGACCAAATTATCAG TAGGGGGCAGGATGAGAAGTGCTGGGGCAGGATGAGAAGTGCTGAGGAGCCAAAGGCACTCAGT GAACCTAGAGGCCAAGGATACATTTTGACATGCTAATAGGCATTTTAGTCATTTGTCCTGCATTT 25 CTTTAGGACAGGCCAAGCTGCCTGGGTCATTGTGAGTCCCAGATAATTCTCTTTGAAATAAAATGT TTTTTAAAGAGAGGGGGAAGGTTGGGGGAGGGTGGTCTGAAGTTAAGAGACTTTGGAGTATTAA GACATTGGATATTTTAGAGAAAATTTTGAACTTTTAAGAAGACTGACCTTTTAAAGTGTTTGAAT ATTATAATTTTTTTAAAAAGACTGTGGGAGCTGGGTGGTGGTATAGGCCCTTTAATCCTAGCACCC AGGAGGCAGAAGCAGCCAGATCTCTGAGTTTGAGACCAGCCTGATCTATAGCATGATTTCCAGGA 30 ATTTTATATTGAGGTGCTGACATTAATATGAAATCTTTGTGAGTGGGCAAGAAAATAAAGACTAA ATTTTTTTTTAGGAATATATCAACCAATTGTTTATTACACAGCATGAACAAAACACAAAAATCAAG CCTTTTCCAGATCTTGCTGACAAGCCTATGGTGTCAAAACTCGGAAACGAGAGGCAGGACCAGGA 35 GTTAAAAGACCAGCGAGGCCTCATGGAGACCTTGTCTCAAGCAGAAATAAACAGGGTTGGTAGCA CACACGAACTCTGAACATCACGAGTGTGCACATACCCACACATGCACCTGTAAAAACAAATCCCC CATCTCCAATGTCTCGTTCTAATCTGTTCTTGTATTAAAAGATAACAAATTTGCCTTTATTA CAAATTTCTCTGCAAACTAGAAAATCTGAAAGATCTATTCCAATTACCTTCTAAATCAAACTACC AGGCTTTGACTCATGCTCAATTCTTGGGTAAATTTGTCATTCGCATGAATCCAAATGTCACACAT 40 CCTATATAATTTAAAGGTTAACAAGTAGAAGAGATGTCCCTAGCACCAAGAAAAGTTTAATCTTA ACAGAAAACAGCTTTCACATCTGCTGTGTGGCACCTTTAACGGCAAGGACGGCGTACAATTCGAA CTGCCCTAATTCTTCTGGTTCTGCCCGATTTTAGCATTCTCAGACGGATCCCATCTCTGATAATG GCAGAAAGTGCAGAGACATCTAAATGGCTCATCTCTGTTCTCATTTCCTTCAAGCTGTCTTTGTC TCTGCTCAATCCGAACAAATCTTCTCATCCTTGCTACAGGTTCTTTCAGCACCGACGACAACAAT 45 GTGTGGACCCTTCTCTGGGAAAGGCGCTTTGAACTTCCCTCATGATTCTTTTCACTTCTGTTCTC CACAGGCTGGTTCTGAACCCGGTGACGAAGGCTGTGATGACGATGATATTTTTGGCCACTTGGCAC TGGGGTTCAGGGTTTTCTCATGACCTTTTACTAGTGTTTCTGGTTGTAGGGTTTCTGAATCA TTGGGGTGAGTCCTCTCCACCTTTCCTCTGAGATCTATCATCTGAGTTTCTGGATACACAACTGG GTCAACTTTCTGTGATGGCTCCATGGCGGTGGGCAGAAGCCTCAAAAAGCCAGCTCCGAACAA 50 AATTGCTAGCTAATCTTTGGAAAGACCTAGACTTTGGCCCCAACTAGCAGACTGAAGTGCTGGAA TAAGGTTAAATCCTTGTGCCACCATGCCTGGACCTAAGCTTTTCATGGCCACTATTCCTCGAGGT CTGGATCAGAAGCTTGTGTATTTCATTTCCGGATTGTCGTTCACTCCAGATTAAAAGTCCAAATG 55 AAAGCAATAGCCATGTAATAATGCCTAGATATAACTCTTCCTTGTTCAGCAGCAAATGCATAAGC

AATAAGCTTAGCTGGGTGGGATCTTCCAAAGCTACTCTGCTCTTTTTTCTTCTTGGACATAGGATT CAGCAACATTCTACTTCTTGATGCCCCTTTATTCTTTGAACCATACATTTTTACTTTTCCTTTCG TAGCTTCTTCCTTTCATCAAAAGATTCTTCATAAGAGTGAAATTTGGGGGTTAGAGAGATGGTTC AGTGGTTAATAGCACTGACTGCTCTTCCAGAGGTCCTGAATTCAATTCCTAGCAACCACATGGTA 5 GCTCATAACCATCTGTAATAGGATCTGATGCCCTCTTTTTGGTGTGTCTGAAGAAGACAGCAACAG AAGGTGAAATTTAACCACACAACAGAATTTATGCCAGGCTTGTTTGAGACTTTTTGTCAAAGCAAT TAATCTAAATCTCTTCACCTTAGCCTCAGGTAGACTCTCTGGACAATGGCAAAAAGCAGCCACAT TCTTCATCAAAATATTACAAGAACGGTCTCTCAGCCACATACTAAAATTCTTCTCTGAAACTTCT 10 AGAGCCAGGCTTCCACAGTTCAAACCACCTTCAGCAACAAAGTCTTCTATATTCCTACGATGATA GCCCTTTAAGCCCCACTTAAAGCATTTCACTGAATTCCAAATCTAAAGTCTCCAAATCTATATTC TTCCAAATAAAAGCATGGTCAGACCTACCTATCACAGCAATATCCCAGTCCCTGGTACCAACCTC TTAATATTTATGTCTATGAGTACACTGTTGCTGTCTTCAGACACACCAGAAGAGGGCATC 15 AGATCTCATTACAAATGGCTGTGAGCCACTACGTAGTTGCTGGGAATTGAACTCAGGACCTCTGG AAGAGCAGCCAGTGCTCTTAACCGCCGAGCCATTTTCTCCAGTCCCAAAGAAACACTTATAAAGG ACAATGTTTTTTTGGTTTTTTTTAAAGGTTTATTTATTTTATGTATATGAGTACACTGTAGCTG TCTTCAGATACACCAGAAGAGGGCATCAGATCTTACTATAGATGGTTGTGAACCACCATGTGGTT 20 CAGTTCTAAAGGACAATGTTTAATCGGGGCTGGCTCACAGGTTCAGAGGTTCAGTCCATTATCAT TGAGACAGGAGCGTGGCAGCATCCAGGCAGGTGTGGGGGCTGAAGGAGCTGAAAGTTCTACCTCTT GATCCAAAGGCAGACCAAAAAAAAAGACTGGCTTACCGGGCTTACCATAAGCAGCTAAGAGGAAGGT CTCAAAGCCCACCCTACAGTGGCATGTTCTCCAACAAGGCCACATCTCCTAATAGTGCCACTCCC CGGGCCATGCATATTCAAGTCGCCACACCCACTGAGCCATCTCCCAACCTGCTCCAGACCATCT 25 CCCCTGCTTTTACCTAAGCTCATTAGGCAGCAATATGCCTCTTATTGTTTGAGCTCAGCATCCTG TTTTTCAAAAGGCTGCTTGTCATCACAGTGGTTTGTTCCACAACTCTCCCAGTTTCTTTGTNAAA ACACCAATGCCTAGAGAGATGCTCTTCTGTACATATCGCATGTGCAGAAGAAAGGGTGCCAGATC CTTTCATGTGGACCNTGTCATGTCTTTACCCACGTAGTCGTCTGCTCTGACTCTTCTCGAGATGC TGANAACTGATTGAGCGTAGGATGCTCTGGGTATGTGCATGGGACAATTTTG

30 SEQ ID NO: 8 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC103122 (11084 - 13244: contig of 2161 bp in length)

CGAAGGACGGTAAGGAGAGAGAGAGGGGAGAGATCAGGACTGAGGGGGAGATATGCACTGAACGGG GGAGTTAGTAACGAGGAAAAGATAGGGAGAAAAAGTGGGAGAAAAAAAGGCCCGGGGAGGGGGAGGGC 35 GGAGTTTTCGGCGAAAGGGCCCGGAGTGTGGATTATCGCGTGGACCAGAACGGGGGAAGGGCCAC TTGAGAAAAAATCATCAAAGCCCCTAAGGAGCATTTGTTTCGGAGTTATACGTATGGATATTTT 40 ATTATATGGGACGAGAGATAAAGAATACTTCTTAAGTAATCCCTTTAAAAAATAATGTCAGGCTGG AGAAATGGTTTCATGGGTAAGCAAGTGTGAGAGATGAGCGCAGACCCCCAGGACCTGTGTAGACT TAATGCAGAGGTGGATGCACGCCTGTAATCTCAGCATGCCTACAGCCAGATAGGAGATGGGGACA GAGAAGTGTGGGGGCCAACTAGCCTGGTGTCTACAGCCTGGTGTCAACAGCAGCCTCCTACCTCA 45 TACTTTACACACATACTCACACTCACACATACATACACATATATACCTGGTCTCCATTAGGCTTC TTTTTGAGGCTTCTGCCCACCACCATGGAGGAGCCATTAGAGAAATCGACCCAGTTGTGGACCCA GAAACTCCTCAGACGAAAGATGAAAAGGACGCATCCGCTGATTCAGAAGTCGTAAGCCAGAAACA CTAGTAAAGGTCATGAAAACGCTAGCCCTGAACCCCAGTGCCAAGCGGTCAGCACATCGTCGCAG CCTCCGTCTCCGGATTCAGAGAAGACCTGTGGAGAACAGAAGTGAAAGAATTTCGAGGGAAGTTC

AAAGCGCTTTACCCAAGAGAAGGGTCCGCACGTTGTTGTCGGTGCTGAGAGATCCTATAGCAAGG GAGTGTGCCATTCAGACTCACTGTGCTTTTCTGCCATTATCAGAGACGGGATCCGTCTGAGAACGC TAAAATCGGGAAGCATTAGGACAGCTTAGATTGTACACTGTCCTTGTGTTAATGATGCCATGCAG CAGACCTGAAAGCTGGCTTTTTGCTTTTTAAGATTAACCTTTTTCCTGGTGCTGGGGACTCTTCTAA CTTGTTAACCTTTAAATTATAGGGTGCGTGATGTTTGGATTCATGTGAATGACTTAAATTTAC CCAAAGAATTGAGAAGGAGTCAAAGCATTCTGTGAATTTTTGAAGCCTCAAGCCCGGGGCCGAGA AACAATGTTAATAGAATTTGGAATAGTTTGGTTTAGAAGGTAATTGGGATAGATCTCTGAATTTT 10 TCAGCCTCCATGTCTTGATCCCAGTCCATCATGAAAGGAAGTCAGGACAGGAACTCAAGTCAGGA CCGTGGAAGTAGCATCTGAAGCAGAGACTTCTGGGATGAAAGCGCTGCTTCCTGACTCGCT CCCCACAAATTGGTCCCTGAGCCTTCTTGTCCACCCTCGGACCCCTTGCCTAGGGTTGGCACCAC CCACAATGGGCTGAGCCTTCCCATGTCAATCACTAATTAAGAAAATGCTGTACAGCGTTGCCTAC TCAAATTGACAACCAGCCAGCCACACAAACANTTAAAAAGATAGAAATAATGTTAGTGNNTC 15 NCATCGAGCAAGAGTC

SEQ ID NO: 9 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC099436 (1 - 21688: contig of 21688 bp in length)

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